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Figure 1A

Identity to SeqID No:1 is indicated by a dot. Gap is indicated by a dash.

* 20 *

Seqid1 : ATGAAAATCCGAGTAAAGTGGCCTATGGTA : 30

Seqid3 : : 30

 40 * 60

Seqid1 : ATGGCGATGGGGCTGGTTTTAAGTGCCTGT : 60

Seqid3 : : 60

 * 80 *

Seqid1 : CAAAGTACACCAATACCGCCTAAAAATAAT : 90

Seqid3 : : 90

 100 * 120

Seqid1 : CCGCAATTGGCACAAATCCGAACCCAAATT : 120

Seqid3 : : 120

 * 140 *

Seqid1 : GCAATCTCGCTACTGGACATGGGTAAGCTT : 150

Seqid3 : : 150

 160 * 180

Seqid1 : GATCAAGCCAAACAGCAGTTGGATGCTGCT : 180

Seqid3 : : 180

* 200 *



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Figure 1B

Seqid1 : TTATCGGCTGATCGGCAGTTTGCACCTGCC : 210
Seqid3 : : 210

220 * 240
Seqid1 : TATCGCACCTTGGCAAAGGTTTATCAAGCT : 240
Seqid3 : : 240

* 260 *
Seqid1 : TCGGAGGATGCCACTCACCAAACCAAAGCT : 270
Seqid3 : : 270

280 * 300
Seqid1 : CAACGCTTGTTTGAAAAGGCGATTGAACTA : 300
Seqid3 : : 300

* 320 *
Seqid1 : AATCCTAAAGATATGCAAAGTTATATGGAT : 330
Seqid3 : : 330

340 * 360
Seqid1 : TATGGATTTTATTTGGTGCAGATGGGGGAC : 360
Seqid3 : : 360

* 380 *
Seqid1 : TTGTCAGGTGCGTTGATTTATTTTGATAAA : 390
Seqid3 : : 390



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400 * 420
Seqid1 : CCTAGCCGAGCCATCGGTTATGAAGGGCGT : 420
Seqid3 : : 420

* 440 *
Seqid1 : GTGGTAGCCATCGAAAATATGGCATATATT : 450
Seqid3 : : 450

460 * 480
Seqid1 : TATTATCATCAATATGAAGCTGCCAAATCA : 480
Seqid3 : : 480

* 500 *
Seqid1 : CCAACAAAAGATGACTATAATAACGCCAAA : 510
Seqid3 : : 510

520 * 540
Seqid1 : TCAGCACTTGAGCGTGCGTTAATTTTCAGGC : 540
Seqid3 : : 540

* 560 *
Seqid1 : ACTCAACATGATGAAATCAAAAAGTCTTAT : 570
Seqid3 : : 570

580 * 600
Seqid1 : GACAAATTATTAAGTGATTATAAATTATTA : 600
Seqid3 : : 600

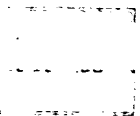


Figure 1D

*

Seqid1 : AGTGATTATAAATAG : 615

Seqid3 :--- : 612

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Figure 2 TECH CENTER 1600/2900

Identity to SeqID No:2 is indicated by a dot. Gap is indicated by a dash.

Seqid2 : MKIRVKWPMVMAMGLVLSACQSTPIPPKNN : 30
Seqid4 : : 30

40 * 60
Seqid2 : PQLAQIRTQIAISLLDMGKLDQAKQQLDAA : 60
Seqid4 : : 60

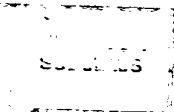
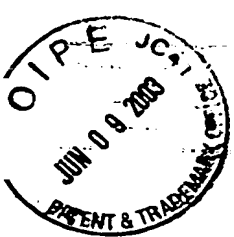
* 80 *
Seqid2 : LSADRQFAPAYRTLAKVYQASEDATHQTKA : 90
Seqid4 : : 90

100 * 120
Seqid2 : QRLF EKAI ELNPKDMQSYMDYGFYLVQM GD : 120
Seqid4 : : 120

* 140 *
Seqid2 : LSGALIYFDKPSRAIGYEGRVVAIENMAYI : 150
Seqid4 : : 150

160 * 180
Seqid2 : YYHQYEAAKSPTKDDYNNAKSALERALISG : 180
Seqid4 : : 180

* 200
Seqid2 : TQHDEIKKSYDKLLSDYKLLSDYK : 204
Seqid4 : : 204



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Fig. 3A

MW BASB121

50 kD →
30 kD →
20 kD →

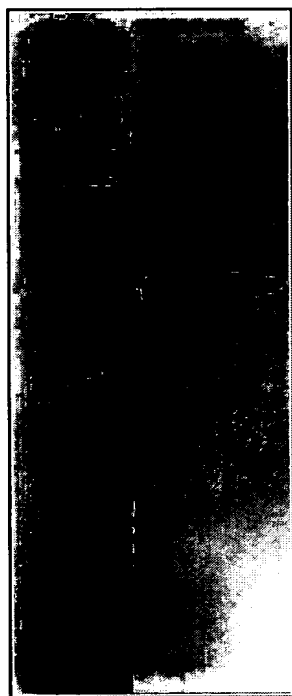


Fig. 3B

MW

BASB121

50 kD →
30 kD →
20 kD →

